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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.	
10/553,509	10/18/2005	Mitsuharu Hirai	TOYA114.010APC	4683	
20995 KNORRE MA	20995 7590 06/04/2007 KNOBBE MARTENS OLSON & BEAR LLP			EXAMINER	
2040 MAIN STREET			WILDER, CYNTHIA B		
FOURTEENTH FLOOR IRVINE, CA 92614			ART UNIT	PAPER NUMBER	
			1637		
			NOTIFICATION DATE	DELIVERY MODE	
			06/04/2007	ELECTRONIC	

Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.

Notice of the Office communication was sent electronically on above-indicated "Notification Date" to the following e-mail address(es):

jcartee@kmob.com eOAPilot@kmob.com

	Application No.	Applicant(s)				
	10/553,509	HIRAI, MITSUHARU				
Office Action Summary	Examiner	Art Unit				
	Cynthia B. Wilder, Ph.D.	1637				
The MAILING DATE of this communication app Period for Reply	pears on the cover sheet with the o	correspondence address				
A SHORTENED STATUTORY PERIOD FOR REPL WHICHEVER IS LONGER, FROM THE MAILING D - Extensions of time may be available under the provisions of 37 CFR 1.1 after SIX (6) MONTHS from the mailing date of this communication. - If NO period for reply is specified above, the maximum statutory period Failure to reply within the set or extended period for reply will, by statute Any reply received by the Office later than three months after the mailin earned patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 136(a). In no event, however, may a reply be time will apply and will expire SIX (6) MONTHS from a, cause the application to become ABANDONE	N. nely filed the mailing date of this communication. D (35 U.S.C. § 133).				
Status						
1)⊠ Responsive to communication(s) filed on 21 N	March 2007.					
	<u> </u>					
3) Since this application is in condition for allowa	Since this application is in condition for allowance except for formal matters, prosecution as to the merits is					
closed in accordance with the practice under t	Ex parte Quayle, 1935 C.D. 11, 4	53 O.G. 213.				
Disposition of Claims						
4) Claim(s) 1-9 is/are pending in the application.						
4a) Of the above claim(s) is/are withdrawn from consideration.						
5) Claim(s) is/are allowed.						
6)⊠ Claim(s) <u>1-9</u> is/are rejected.						
7) Claim(s) is/are objected to.						
8) Claim(s) are subject to restriction and/o	or election requirement.					
Application Papers						
9) The specification is objected to by the Examine	er.					
10) The drawing(s) filed on is/are: a) acc		Examiner.				
Applicant may not request that any objection to the						
Replacement drawing sheet(s) including the correct	tion is required if the drawing(s) is ob	jected to. See 37 CFR 1.121(d).				
11) ☐ The oath or declaration is objected to by the E	xaminer. Note the attached Office	Action or form PTO-152.				
Priority under 35 U.S.C. § 119	•					
12) ☐ Acknowledgment is made of a claim for foreign a) ☐ All b) ☐ Some * c) ☐ None of:	n priority under 35 U.S.C. § 119(a)-(d) or (f).				
1. Certified copies of the priority document	1. Certified copies of the priority documents have been received.					
2. Certified copies of the priority documents have been received in Application No						
3. Copies of the certified copies of the price	•	ed in this National Stage				
application from the International Burea	· · · · · · · · · · · · · · · · · · ·	_				
* See the attached detailed Office action for a list	t of the certified copies not receive	ea.				
Attachment(s)	,					
1) Notice of References Cited (PTO-892)	4) Interview Summary					
2) Notice of Draftsperson's Patent Drawing Review (PTO-948) 3) Information Disclosure Statement(s) (PTO/SB/08)	Paper No(s)/Mail D 5) Notice of Informal F					
Paper No(s)/Mail Date	6) Other:					

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FINAL ACTION

1. Applicant's amendment filed 3/21/2007 is acknowledged and has been entered. Claims 1-9 are pending and discussed in this Office action. All of the arguments have been thoroughly reviewed and considered but are not found persuasive for the reasons discussed below. Any rejection not reiterated in this action has been withdrawn as being obviated by the amendment of the claims.

This action is made FINAL.

2. The text of those sections of Title 35, U.S. Code not included in this action can be found in a prior Office action.

Previous Rejections

- 3. The objection to the priority documents is withdrawn in view of Applicant's arguments. The prior art rejection under 35 USC 103(a) is maintained and discussed below.
- 4. <u>Issue I. Claims 1-2, and 7-9 are rejected under 35 USC 103(A) as being unpatentable over Shuldiner et al in view of Hiratsuka et al.</u>

Applicant's traversal

(I) Applicant traverses the rejection on the following grounds: Applicant states that the nucleic acid probes of the claimed invention have a nucleotide sequence starting at nucleotide 183 of SEQ ID NO: 1 and a length of 8 to 30 nucleotides or a nucleotide sequence ending at nucleotide 196 of SEQ ID NO: 2 and a length of 7 to 30 nucleotides. Applicant states that this feature is not taught by either of the cited

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references. Applicant states that neither Shuldiner nor Hiratsuka et al teach probes having the specific endpoint as claimed. Applicant states that among the many cytosines that could be labeled for detection of $\beta 3$ Adrenaline receptor mutations, the cytosines at 183 of SEQ ID NO: 2 and 196 of SEQ ID NO: 2 are critical for detecting the Trp64Arg mutation by Tm analysis. Applicant states that when probes as claimed are used, which have cytosine at the 5' end (position 183) or 3' (position 196), changes in fluorescence intensity that could be analyzed by Tm analysis were observed. Applicant states however, when probes having cytosine other than cytosine 183 or 196 at the 5' or 3' end, respectively, were used, changes in fluorescence intensity were not observed. Applicant states that Shuldiner, et al discloses only a part of the nucleotide sequence having the β3AR mutation and fails to teach the criticality of nucleotides 183 and 196. Applicant states that Shuldiner et al do not teach or suggest that the cytosine at position 183 and 196 corresponds to the 5' end and 3' end of the probes respectively. Applicant states that this deficiency is not corrected by Hiratsuka et al. Applicant states that Hiratsuka et al discloses a probe labeled with fluorescence tag and a method for detecting single nucleotide polymorphisms by using the probe. Applicant states that the claims do not teach the specific positions in the probes as claimed. Applicant states that accordingly, one of ordinary skill in the art would not have a reasonable expectation of success in achieving the claimed invention by combining Shuldiner et al and Hiratsuka et al. Applicant states that the importance of the cytosines at positions 183 and 196 for the 5' and 3' ends of the probe, respectively, could not have been predicted based upon the cited references. Applicant asserts that the claims 2-9 are

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believed to patentable at least because they include all of the limitations of claim 1, which Applicant asserts is patentable for the reasons given above. Applicant request reconsideration and withdrawn of the rejections.

Examiner's Response

(ii) All of the arguments have been thoroughly reviewed and considered but are not found persuasive for the reasons that follow: In regards to Applicant arguments that the references does not teach the instant invention or the criticality of the cytosine at position 183 of SEQ ID NO: 1 and 196 of SEQ ID NO: 2, the Examiner respectfully Firstly, the courts have established that during patent examination the pending claims must be interpreted as broadly as their terms reasonably allow (In re Zletz, 893 F.2d 321-22, 13 USPQ2d 1320, 1322 (Fed. Cir. 1989). In this case, the claims as broadly written do not require that a cytosine or any other specific base be position at nucleotide 183 of SEQ ID NO: 1. The claims only require that the probe sequence start at position 183 of SEQ ID NO: 1 and have a length of 8 to 30 nucleotides and the 5' end labeled with a fluorescent dye. Alternatively, the claims require that the nucleotide probe end at the nucleotide number 196 in the nucleotide sequence of SEQ ID NO: 2 and having a length of 7 to 30 nucleotides and wherein the 3' end of the probe is labeled with a fluorescent dye. The claims, especially claim 2. suggest that the probe comprising the sequences as recited in SEQ ID NOS: 8-12 corresponds to the probe as broadly recited in the claim 1.

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Thus, contrary to Applicant's arguments, the prior art of Shuldiner does teach the instant invention as claimed. Shuldiner et al provides a probe nucleotide sequence that is 100% identical to the sequences of SEQ ID NOS: 11 and 12 (SEE SEQ ID NO: 7 and Appendix A, B and C at this end of this Office action). An alignment of this sequence with the sequence of SEQ ID NO: 1 shows that this sequence begins at nucleotide 183 of SEQ ID NO: 1. Additionally, while the claim does not require that a cytosine be present at position 183, the sequence of Shuldiner provides this teaching as well as a cytosine is present at the first position or 5' end of the probe of Shuldiner et al (see SEQ ID NO: 7 at col. 25 or Appendix A). Further, Shuldiner teaches wherein the probe meets the length limitation of the probe of claim 1 (Shuldiner teach wherein the probes are 17 nucleotides in length) and additionally teaches wherein the probe is used to detect the Trp64Arg mutation (see columns 11, 12, 17 and 18). The secondary reference of Hiratsuka provides the limitations not found in the patent of Shuldiner and provides motivation for combining the teachings. Applicant's arguments are not sufficient to overcome the prior art rejections. Accordingly, the rejections are maintained.

5 <u>Issue II. Claims 3-6 are rejected under 35 USC 103(A) as being unpatentable over Hiratsuka et al in view of Shuldiner et al as previously applied above.</u>

Applicant's traversal and Examiner's Response

(I) Applicant traverses the rejection on the same grounds presented above at number 4. The Examiner in response asserts that Applicant's arguments are not

sufficient to overcome the prior art rejections under 35 USC 103(a) and in conclusion maintains the rejections for the reasons discussed above.

Conclusion

6. No claims are allowed. THIS ACTION IS MADE FINAL. Applicant is reminded of the extension of time policy as set forth in 37 CFR 1.136(a).

A shortened statutory period for reply to this final action is set to expire THREE MONTHS from the mailing date of this action. In the event a first reply is filed within TWO MONTHS of the mailing date of this final action and the advisory action is not mailed until after the end of the THREE-MONTH shortened statutory period, then the shortened statutory period will expire on the date the advisory action is mailed, and any extension fee pursuant to 37 CFR 1.136(a) will be calculated from the mailing date of the advisory action. In no event, however, will the statutory period for reply expire later than SIX MONTHS from the mailing date of this final action.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Cynthia B. Wilder, Ph.D. whose telephone number is (571) 272-0791. The examiner can normally be reached on a flexible schedule.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Gary Benzion can be reached on (571) 272-0782. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

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Business Center (EBC) at 866-217-9197 (toll-free). If you would like assistance from a

USPTO Customer Service Representative or access to the automated information

system, call 800-786-9199 (IN USA OR CANADA) or 571-272-1000.

Cynthia B. Wilder, Ph.D. Patent Examiner

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PRIMARY EXAMINER

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5/24/07

(x i) SEQUENCE DESCRIPTION: SEQ ID NO:12:

26

(i i) MOLECULE TYPE: DNA (genomic) (x i) SEQUENCE DESCRIPTION: SEQ ID NO:7: Instant invention. CATCGCCTGG ACTCCGA (2) INFORMATION FOR SEQ ID NO:8: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: mucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (i i) MOLECULE TYPE: DNA (genomic) (x i) SEQUENCE DESCRIPTION: SEQ ID NO:8: CATCGCCCGG ACTCCGA 1 7 (2) INFORMATION FOR SEQ ID NO:9: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (i i) MOLECULE TYPE: DNA (genomic) (x i) SEQUENCE DESCRIPTION: SEQ ID NO:9: GAAAGGGGAC AGATCTCACC AA 2 2 (2) INFORMATION FOR SEQ ID NO:10: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: mucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (i i) MOLECULE TYPE: DNA (genomic) (x i) SEQUENCE DESCRIPTION: SEQ ID NO:10: TTCCTCTGCC ACCATCTGT 19 (2) INFORMATION FOR SEQ ID NO:11: (i) SEQUENCE CHARACTERISTICS: (A) LENOTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (i i) MOLECULE TYPE: DNA (genomic) (x i) SEQUENCE DESCRIPTION: SEQ ID NO:11: CCAGGGAGTG CTATGCTGAG 20 (2) INFORMATION FOR SBQ ID NO:12: (i) SEQUENCE CHARACTERISTICS: (A) LENOTH: 24 base pairs (B) TYPE: anclaic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (i i) MOLECULE TYPE: DNA (genomic)

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ppendix B
<!--StartFragment-->US-08-446-530-7
; Sequence 7, Application US/08446530
;.Patent No. 5766851
  GENERAL INFORMATION:
    APPLICANT: Shuldiner, Alan R.
    APPLICANT: Walston, Jeremy
    APPLICANT: Silver, Kristi
    TITLE OF INVENTION: SUSCEPTIBILITY GENE FOR OBESITY AND TYPE
    TITLE OF INVENTION: II DIABETES MELLITUS
    NUMBER OF SEQUENCES: 28
    CORRESPONDENCE ADDRESS:
      ADDRESSEE: Fish & Richardson P.C.
      STREET: 4225 Executive Square
      CITY: La Jolla
      STATE: CA
COUNTRY: USA
      ZIP: 92037
    COMPUTER READABLE FORM:
      MEDIUM TYPE: Floppy disk
      COMPUTER: IBM PC compatible
      OPERATING SYSTEM: PC-DOS/MS-DOS
      SOFTWARE: PatentIn Release #1.0, Version #1.30
    CURRENT APPLICATION DATA:
      APPLICATION NUMBER: US/08/446,530
      FILING DATE: 19-MAY-1995
      CLASSIFICATION: 435
    ATTORNEY/AGENT INFORMATION:
      NAME: Haile, Lisa A.
      REGISTRATION NUMBER: 38,347
      REFERENCE/DOCKET NUMBER: 07265/048001
    TELECOMMUNICATION INFORMATION:
      TELEPHONE: 619/678-5070
      TELEFAX: 619/678-5070
   INFORMATION FOR SEQ ID NO: 7:
    SEQUENCE CHARACTERISTICS:
      LENGTH: 17 base pairs
      TYPE: nucleic acid
      STRANDEDNESS: single
      TOPOLOGY: linear
    MOLECULE TYPE: DNA (genomic)
                                         SED ID NO! 11
US-08-446-530-7
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 Query Match
                        100.0%; Pred. No. 30;
 Best Local Similarity
 Matches 16; Conservative 0; Mismatches 0; Indels
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                                                                          0;
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Qу
             1 CATCGCCTGGACTCCG 16
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<!--EndFragment-->

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<!--StartFragment-->RESULT 2
                                         Appendix C
US--08-446-530-7
;.Sequence 7, Application US/08446530
; Patent No. 5766851
  *GENERAL INFORMATION:
    APPLICANT: Shuldiner, Alan R.
    APPLICANT: Walston, Jeremy
    APPLICANT: Silver, Kristi
    TITLE OF INVENTION: SUSCEPTIBILITY GENE FOR OBESITY AND TYPE
    TITLE OF INVENTION: II DIABETES MELLITUS NUMBER OF SEQUENCES: 28
    CORRESPONDENCE ADDRESS:
      ADDRESSEE: Fish & Richardson P.C.
      STREET: 4225 Executive Square
      CITY: La Jolla
      STATE: CA
      COUNTRY: USA
      ZIP: 92037
    COMPUTER READABLE FORM:
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      COMPUTER: IBM PC compatible
      OPERATING SYSTEM: PC-DOS/MS-DOS
      SOFTWARE: PatentIn Release #1.0, Version #1.30
    CURRENT APPLICATION DATA:
      APPLICATION NUMBER: US/08/446,530
      FILING DATE: 19-MAY-1995
      CLASSIFICATION: 435
    ATTORNEY/AGENT INFORMATION:
      NAME: Haile, Lisa A.
      REGISTRATION NUMBER: 38,347
      REFERENCE/DOCKET NUMBER: 07265/048001
    TELECOMMUNICATION INFORMATION:
      TELEPHONE: 619/678-5070
      TELEFAX: 619/678-5070
   INFORMATION FOR SEQ ID NO:
    SEQUENCE CHARACTERISTICS:
      LENGTH: 17 base pairs
      TYPE: nucleic acid
      STRANDEDNESS: single
      TOPOLOGY: linear
    MOLECULE TYPE: DNA (genomic)
                               sto SEQ IDNO:12
US-08-446-530-7
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  Query Match
  Best Local Similarity 100.0%; Pred. No. 54;
                                                               0; Gaps
  Matches 15; Conservative 0; Mismatches
                                                 0; Indels
           1 CATCGCCTGGACTCC 15
Qу
             1 CATCGCCTGGACTCC 15
```

<!--EndFragment-->